A Preclinical Evaluation of Neural Stem Cell–Based Cell Carrier for Targeted Antiglioma Oncolytic Virotherapy

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Background Oncolytic adenoviral virotherapy (OV) is a highly promising approach for the treatment of glioblastoma multiforme (GBM). In practice, however, the approach is limited by poor viral distribution and spread throughout the tumor mass.

Methods To enhance viral delivery, replication, and spread, we used a US Food and Drug Administration–approved neural stem cell line (NSC), HB1.F3.CD, which is currently employed in human clinical trials. HB1.F3.CD cells were loaded with an oncolytic adenovirus, CRAd-Survivin-pk7, and mice bearing various human-derived GBMs were assessed with regard to NSC migration, viral replication, and therapeutic efficacy. Survival curves were evaluated with Kaplan–Meier methods. All statistical tests were two-sided.

Results Antiglioma activity of OV-loaded HB1.F3.CD cells was effective against clinically relevant human-derived glioma models as well as a glioma stem cell–enriched xenograft model. Median survival was prolonged by 34% to 50% compared with mice treated with OV alone (GBM43FL model median survival = 19.5 days, OV alone vs NSC + OV, hazard ratio of survival = 2.26, 95% confidence interval [CI] = 1.21 to 12.23, P = .02; GBM12 model median survival = 43.5 days, OV alone vs NSC + OV, hazard ratio of survival = 2.53, 95% CI = 1.21 to 10.38, P = .02). OV-loaded HB1.F3.CD cells were shown to effectively migrate to the contralateral hemisphere and hand off the therapeutic payload of OV to targeted glioma cells. In vivo distribution and migratory kinetics of the OV-loaded HB1.F3.CD cells were successfully monitored in real time by magnetic resonance imaging. OV-loaded NSCs retained their differentiation fate and were nontumorigenic in vivo.

Conclusions HB1.F3.CD NSCs loaded with CRAd-Survivin-pk7 overcome major limitations of OV in vivo and warrant translation in a phase I human clinical trial for patients with GBM.

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Neural stem cells (NSCs) are defined as multipotent progenitor cells that originate from the developing and adult central nervous system (1). NSCs display intrinsic tumor tropism that can be exploited for targeted anticancer drug delivery to invasive and metastatic cancer (2,3). In theory, the tumor-homing property of NSCs offers a substantial advantage over other targeted therapies, such as antibody-directed drug delivery, because of their ability to detect various cues generated by satellite tumor foci and respond to such cues by extravasating through complex tissue microenvironments and migrating to distant diseased areas (4). Glioblastoma multiforme (GBM) is the most common and aggressive primary central nervous system tumor in adults and is characterized by its propensity to infiltrate throughout the brain and cause relapses in patients because of the existence of an aberrant chemo- and radio-resistant glioma stem cell (GSC) population (5). Thus, a true cure for this formidable disease cannot arise from the application of traditional antineoplastic principles; it requires a dynamic agent capable of targeting scattered disease lesions as well as eliminating the tumor-initiating cancer stem cells effectively with minimal disruption of the existing delicate neural architecture (6).

Based on this, our lab has extensively evaluated NSCs as a cellular vehicle for the targeted delivery of glioma tropic oncolytic adenoviral virotherapy (OV) CRAd-S-pk7 (7–9). We have proposed that by combining NSCs’ unique tumor tropism with the OV’s ability to target chemo- and radio-resistant GSCs (6,10) we may overcome the deficiencies inherent to each approach deployed in isolation and can effectively target GBM. As a proof-of-principle we have previously demonstrated that 1) NSCs can be used as cellular vehicles for the in vivo delivery of an OV to intracranial gliomas (7); 2) intratumoral delivery of NSCs loaded with the CRAd-S-pk7, a glioma-tropic OV regulated by the tumor-specific survivin promoter (7,11), increased median survival by 50% compared with mice treated with OV alone in an orthotopic xenograft model of human glioma (8); and 3) NSCs...
demonstrated superior therapeutic efficacy when compared with mesenchymal stem cells as a cell carrier for OV in the context of intracranial gliomas (9). Because these previously published results argue in favor of using NSCs as targeted cellular delivery vehicles for antiglioma oncolytic virotherapy, we conducted the following critical translational studies to justify its application in a phase I clinical trial for patients with GBM: 1) identified an optimal NSC-based cell carrier for antiglioma oncolytic virotherapy; 2) tested the selected NSC-based cell carrier in several diverse and clinically relevant glioma xenograft models; 3) developed a noninvasive imaging protocol to monitor in vivo distribution and migratory activity of NSC-based cell carriers in real time; 4) examined the capacity for the NSC-based cell carrier to deliver antiglioma OV to a distant tumor burden in a glioma xenograft model; and 5) evaluated the therapeutic efficacy of NSC-based oncolytic virotherapy in a distance-delivery glioma xenograft model.

In this report, we provide a detailed evaluation of two immortalized NSC lines as cell carriers for targeted antiglioma therapy. Our results indicate that HB1.F3.CD, a US Food and Drug Administration–approved NSC line for human clinical trials (NCT01172964) is the most suitable NSC cell carrier for the future application of cell-based OV delivery in the clinical setting. We found that HB1.F3.CD cells can effectively hand off the viral therapeutic payload to distant tumor sites and substantially prolong median survival in diverse orthotopic models of human glioma. Thus, data presented in this study solidify the notion that NSCs can be used as cell carriers for the targeted delivery of antiglioma oncolytic viruses and serve as the foundation for an investigational new drug application for a human clinical trial involving newly diagnosed and recurrent patients with malignant gliomas.

Methods

Cell Culture and Establishment of Fluorescent-Labeled Cell Lines

The U87MG, U118MG, U251, and A172 human glioma cell lines were purchased from the American Type Culture Collection (Manassas, VA) and maintained according to vendor recommendations. Human primary brain tumor specimens (GBM43 and GBM12) were obtained from Dr David James (University of California at San Francisco) in accordance to a protocol approved by the institutional review board at UCSC. Tumor specimens were confirmed as World Health Organization grade IV malignant glioma by an attending neuropathologist. Human ReNcells (NSCs) were obtained from Millipore (Temecula, CA) and maintained according to the manufacturer’s protocol. Briefly, these NSCs were isolated from the cortical region of 14-week-old fetal tissue and immortalized by retroviral transduction and insertion of the c-myc gene. Cells were characterized according to the expression of nestin, SOX-2, CD133, and CD44 (data not shown) stem cell markers. Further details are available in the Supplementary Methods (available online).

Viral Vector

The replication competent adenoviral vector CRAd-S-pk7 contains the wild-type adenovirus replication protein E1A under the control of human survivin promoter. This vector has been created by homologous recombination using a shuttle plasmid containing the human survivin promoter upstream to the viral E1A gene. Shuttle plasmids containing these regions were further homologously recombined into adenoviral plasmids modified to contain a poly-lysine (pk7) incorporation into the C-terminus of the wild-type fiber protein (7,11).

Analysis of Viral Replication

To detect the level of viral replication by quantitative polymerase chain reaction (PCR), NSCs were plated at a density of 2.5 × 10^4 cells/well in 24-well plastic tissue culture dishes. The next day, cells were infected with indicated IU/cell of CRAd-S-pk7. After a 1-hour incubation period, virus-containing media was removed, cells were washed with 1X phosphate-buffered saline (PBS), and a fresh portion of complete growth media was added. Infected cells were harvested at indicated time points. Total DNA was extracted from infected cells using a DNeasy Tissue Kit (Qiagen, Hilden, Germany) according to the manufacturer’s protocol. Gene expression was quantified by real-time quantitative PCR using SYBR Green PCR Master Mix (Applied Biosystems, Foster City, CA) and primers recognizing the viral E1A gene (7). In this process, DNA was amplified using an Opticon 2 system (Bio-Rad, Foster City, CA) and was detected by measuring the binding of the fluorescent dye SYBR green. Each sample was run in triplicates. Results were presented as the average number of E1A copies per nanogram of DNA.

We used the Adeno-X Rapid Titer Kit (Clontech, Mountain View, CA) according to the manufacturer’s protocol to titrate the levels of infectious viral progeny. Briefly, infected cells and media from each group were collected and subsequently subjected to three cycles of freezing and thawing. As a consequence, infectious progenies were released from infected cells. Cell lysates were then incubated with adherent HEK293 cells in serial 10-fold dilution. Forty-eight hours later, the amount of IU was calculated using the Adeno-X Rapid Titer Kit according to vendor recommendations. The titration units (IU) used by this protocol are similar to plaque-forming units.

Evaluation of NSC Migration and Viral Delivery In Vitro

To analyze the migratory capacity and OV delivery characteristics of HB1.F3-CD and ReNcells in vitro, we used a similar system to that described earlier (8), with a slight modification. Further details are available in the Supplementary Methods (available online).

Animal Studies

U87MG glioma cells were implanted by cranial guide screw. Briefly, mice were anesthetized with a ketamine/xylazine mixture (115/17 mg/kg), and a burr hole was made. Stereotactic injection was carried out by using a 10 μL Hamilton syringe (Hamilton, Reno, NV) with a 30-gauge needle, which was inserted through the burr hole to a depth of 3 mm mounted on a mice-exclusive stereotactic apparatus (Harvard Apparatus, Holliston, MA). Male athymic/nude mice were obtained from Charles River Laboratory (Wilmington, MA). Mice were cared for according to a study-specific animal protocol approved by the University of Chicago Institutional Animal Care and Use Committee. For detailed description of the methods used for all the in vivo experiments, please see the Supplementary Methods (available online).
Statistical Analysis
All statistical analyses were performed using Graphpad Prism 4 (GraphPad Software, San Diego CA). Data represent the results for assays performed in triplicate and repeated at least three different times, and error bars represent 95% confidence intervals (CIs). For continuous variables, comparisons between two groups were evaluated by statistical significance of difference of means in independent sample sets and determined using the Student t-test; comparisons between more than two treatment groups were made using one-way analysis of variance or Kruskal–Wallis with Dunnett post hoc test. Survival curves were generated by the Kaplan–Meier method, and the log-rank test was used to compare the distributions of survival times. All reported P values were two-sided and were considered to be statistically significant at less than .05.

Results
Evaluation of CRAd-S-pk7 Progeny and Hand-Off From NSCs to Glioma Cells
The initial characterization of both the HB1.F3.CD and ReNcell NSC lines and optimization of OV loading dose for the carrier cells is described in the Supplementary Results section and Supplementary Figure 1 (available online). Next, we examined in both NSC lines the capacity of released viral progeny to lyse targeted tumor cells. Both NSC lines were infected with 50 IU of OV per cell, the optimal OV loading dose according to our previous experiments [Supplementary Figure 1 and ref. (8)] and placed in the upper chambers of a transwell plate, whereas glioma cells were cultured in the lower chamber at the NSC/glioma cell ratios of 1:2, 1:10, 1:50, and 1:100. Viral E1A copies in the target glioma cells were measured by quantitative PCR from total isolated DNA. At the NSC/glioma cell ratio of 1:50 and 1:100, the recovered E1A copies from glioma cells were lower by approximately 1.3 (95% CI = 0.84 to 1.81; P = .002) and 1.4 (95% CI = 0.47 to 2.25; P = .02) orders of magnitude, respectively, in the cocultures containing ReNcells as compared with HB1.F3.CD cells (Figure 1A). Moreover, CRAd-S-pk7–loaded HB1.F3.CD cells demonstrated statistically significantly more killing of the targeted glioma cells than ReNcells at the NSC/glioma cell ratio of 1:10 (ReN vs HB1.F3.CD = 39.9% vs 18.7%, difference = 21.2%, 95% CI = 14.4% to 28.1%, P < .001), 1:50 (ReN vs HB1.F3.CD = 57.5% vs. 39.9%, difference = 17.6%, 95% CI = 3.2% to 32.0%, P = .02), and 1:100 (ReN vs HB1.F3.CD = 80.1% vs 53.6%, difference = 26.4%, 95% CI = 5.8% to 47.0%, P = .02) (Figure 1B).

Next, to evaluate the therapeutic efficacy of different NSC-based antiglioma virotherapy in vivo, 2.5 × 10^4 cells of U87MG were implanted into the right hemisphere of athymic nude (nu/nu) mice, grown to 60% confluence, and infected with 50 IU per cell of CRAd-S-pk7. Three days after glioma establishment, both NSC lines were infected with CRAd-S-pk7 in a volume of 2.5 × 10^5 IU per mouse 2 to 3 mm away from the original tumor site. Two additional groups of mice received either 2.5 × 10^5 IU of CRAd-S-pk7 alone or phosphate-buffered saline (PBS) in an identical volume and location in the brain. Survival curves were obtained by the Kaplan–Meier method, and overall survival time was compared between groups using log-rank test. All statistical tests were two-sided.

Figure 1. Permissiveness of neural stem cell (NSC) lines for adenovirus replication and efficacy in vitro/vivo. A) A transwell chamber assay was used to measure the viral progeny released from NSCs and subsequently capable of infecting glioma cells. HB1.F3.CD and ReNcell were infected with 50 IU per cell of CRAd-S-pk7 in the upper chambers of the transwell assay in the following ratio to glioma cells: 1:2, 1:10, 1:50, and 1:100. Glioma cells were placed on the bottom chambers of the transwell assay, and cells were harvested and infectivity was measured for assays performed in triplicate and repeated at least three different times, and error bars represent 95% confidence intervals. Student t test was used. *P < .05; **P < .01. B) Cytotoxicity by trypan blue exclusion 96 hours after coculture. Bars represent means from five independent experiments; error bars refer to 95% confidence intervals. Student t test was used. *P < .05; **P < .01. C) CRAd-S-pk7 virus-loaded NSCs inhibit xenograft growth and prolong survival of mice with orthotopic glioblastoma. We stereotactically injected 2.5 × 10^7 U87MG cells into the right hemisphere of the brains of athymic nude mice (n = 8 per group). Three days after glioma establishment, both NSC lines were infected with 50 IU per cell of CRAd-S-pk7. Separate groups of mice received an injection of either 5 × 10^5 HB1.F3.CD or ReNcell cells loaded with CRAd-S-pk7 in a volume of 2.5 μL per mouse 2 to 3 mm away from the original tumor site. Two additional groups of mice received either 2.5 × 10^5 IU of CRAd-S-pk7 alone or phosphate-buffered saline (PBS) in an identical volume and location in the brain. Survival curves were obtained by the Kaplan–Meier method, and overall survival time was compared between groups using log-rank test. All statistical tests were two-sided.
three days after implantation of the glioma xenograft, a single injection of CRAd-S-pk7 (50 IU of pk7 per NSC x 5 x 10⁶ NSCs per mouse = 2.5 x 10⁹ IU of OV total) or 5 x 10⁷ NSCs loaded with 50 IU per cell of CRAd-S-pk7 was administered ipsilaterally 2 to 3 mm away from the original tumor implantation site. The mice were monitored for survival. The median survival for the PBS-treated control group was 64 days compared with 70.5 days for the CRAd-S-pk7 group (P = .10), 79.5 days for the OV-loaded ReNcell group (P = .06), and 108.5 days for the OV-loaded HB1.F3.CD group (P = .005) (Figure 1C). Thus, the group of animals bearing glioma xenografts treated with CRAd-S-pk7-loaded HB1 survived 29 days longer than the group treated with CRAd-S-pk7-loaded ReNcell. Based on this, we concluded that the HB1.F3.CD cell line functions more effectively as a cell carrier for the OV CRAd-S-pk7 both in vitro and in vivo and used it as our OV carrier cell in the remainder of the studies.

**Evaluation of HB1.F3.CD as a Cell Carrier in Multiple Clinically Relevant GBM Patient-Derived Glioma Xenograft Models**

Next, we tested our OV-NSC system in two patient-derived orthotopic models: GBM43FL and GBM12FL. To maintain their tropism for tumor as detected by MRI in vivo and that MRI contrast, no obvious signal changes were observed along the corpus callosum in the animal brain without tumor (Figure 2A, image rows with tumor). In contrast, Prussian blue staining of the same mouse brain with tumor as shown in Figure 2C, the hypointense stream directed toward the side of the implanted U87 tumor extended gradually toward the side of the implanted U87 tumor mass (Figure 2D). The hazard ratio of survival was 2.53 (95% CI = 1.21 to 10.38, P = .02). In the GBM12FL model, an identical injection protocol was used as GBM43FL line for orthotopic models: GBM43FL and GBM12FL. T o maintain their tropism for tumor as detected by MRI in vivo and that MRI contrast, no obvious signal changes were observed along the corpus callosum in the animal brain without tumor (Figure 2A, image rows with tumor). In contrast, Prussian blue staining of the same mouse brain with tumor as shown in Figure 2C, the hypointense stream directed toward the side of the implanted U87 tumor extended gradually toward the side of the implanted U87 tumor mass (Figure 2D). The hazard ratio of survival was 2.53 (95% CI = 1.21 to 10.38, P = .02). In the GBM12FL model, an identical injection protocol was used as GBM43FL line for orthotopic models: GBM43FL and GBM12FL. T o maintain their tropism for tumor as detected by MRI in vivo and that MRI contrast, no obvious signal changes were observed along the corpus callosum in the animal brain without tumor (Figure 2A, image rows with tumor). In contrast, Prussian blue staining of the same mouse brain with tumor as shown in Figure 2C, the hypointense stream directed toward the side of the implanted U87 tumor extended gradually toward the side of the implanted U87 tumor mass (Figure 2D). The hazard ratio of survival was 2.53 (95% CI = 1.21 to 10.38, P = .02).
The in vivo differentiation status of CRAd-S-pk7–loaded HB1.F3.CD cells is an important issue to consider for both safety (nononcogenic) and tumor-homing properties of loaded NSCs (14). To examine the differentiation status of transplanted HB1.F3.CD cells loaded with OV in vivo, HB1.F3.CD-GFP+ cells infected with OV were implanted in the contralateral hemisphere of nude mouse brains containing U87 glioma xenografts established 10 days earlier. At 24 and 72 hours after NSC implantation, the mice were killed and their brains were subjected to immunohistochemical analysis. Within 24 hours of implantation, the GFP-tagged HB1.F3.CD cells began to cross the midline of the brain and migrate toward the implanted tumor.
Figure 3. (A) GBM43FL glioma cells were cultured and exposed to a total of 10 Gy radiotherapy (XRT) treatment (6 days × 2 Gy), temozolomide (50 µM), or cocultured with HB1.F3.CD-GFP+ cells loaded with CRAd-S-pk7 (50 IU per cell). After 72 hours of incubation, cells were collected and stained for glioma stem cell (GSC) markers CD133 and CD16 and subjected to fluorescent-activated cell sorting (FACS) analysis. Representative FACS plots show the percentage of CD15+, CD133+, or CD15−CD133− GBM43FL cells. (B) The percentage of positive populations of GSCs in the three treatment groups. All three populations of GSCs were statistically significantly reduced in oncolytic adenoviral virotherapy (OV)–loaded NSC treatment group compared with the XRT or chemotherapy (TMZ) treatment groups (P < .001), compared using Student t test. Bars represent means from three independent experiments; error bars refer to 95% confidence intervals. (C) The OV-loaded HB1.F3.CD therapy was tested in vivo. GBM43FL cells were FACS sorted, and 5 × 10^6 CD133+ cells were intracranially implanted in the brains of nude mice (n = 7 per group). Three days after tumor implantation, mice were treated either with phosphate-buffered saline (PBS), 5 × 10^6 HB1.F3.CD cells, 5 × 10^6 HB1.F3.CD cells loaded with 50 IU per cell of CRAd-S-pk7, or 2.5 × 10^7 IU of CRAd-S-pk7 intratumorally. P1 represents P value 1 in this table where PBS treated group served as reference and P2 represents P value 2 where CRAd-S-pk7+HB1.F3.CD serve as reference. Survival curves were obtained by the Kaplan–Meier method, and overall survival time was compared between groups using the log-rank test. All statistical tests were two-sided. ***P < .001; **P < .01; *P < .05.

In the contralateral hemisphere. Figure 5A shows the hematoxylin and eosin staining of the migratory path of implanted HB1.F3.CD OV-loaded cells, which is validated by the immunofluorescence staining by both 4′,6-diamidino-2-phenylindole (DAPI) (Figure 5, B1) and GFP (Figure 5, B2) antibodies on the same animal brain. Next, to evaluate the stemness of implanted HB1.F3.CD cells loaded with CRAd-S-pk7, we counterstained the slides with an antihuman-specific nestin antibody and observed that the majority of the GFP+ HB1.F3.CD cells were also nestin+ (Figure 5, B3, B3-1, and B3-2). When we examined the implanted xenograft tumor in the contralateral hemisphere 72 hours after NSC implantation, we observed the presence of GFP+ cells (Figure 5, C2) inside human CD4+ tumor foci (Figure 5, C1). Additionally, the HB1.F3.CD cells that migrated and reached the tumor site (white dotted area) also stained positive for human nestin (Figure 5, C3 and C4). Furthermore, as shown in Supplementary Figure 2 (available online), even though 24 hours after OV infection of HB1.F3.CD cells in vitro the NSC markers Sox2 and Oct4 as well as the differentiation markers Galc (oligodendrocyte marker), GFAP (astrocyte), and beta-III tubulin (neuronal) mRNA levels decreased, by 72 hours gene expression normalized and was non-statistically significantly different from the uninfected control HB1.F3.CD cells (Supplementary Figure 3, available online). Most important, OV-loaded HB1.F3.CD cells implanted intracranially into nude mice did not show any signs that implanted NSCs became tumorigenic as demonstrated by complete animal survival and immunohistochemistry (data not shown). Taken together, these data indicate that implanted HB1.F3.CD cells loaded with CRAd-S-pk7 retained their NSC differentiation fate and displayed substantial pathotropism in a mouse model of glioma.

In Vivo Hand-off and Expansion of CRAd-S-pk7 at Distant Tumor Sites by HB1.F3.CD Cell Carrier

To investigate the hand-off and amplification of CRAd-S-pk7 at the targeted tumor site, we implanted OV-loaded HB1.F3.CD cells in the contralateral hemisphere of nude mice bearing U87 xenograft tumors (10 days after tumor implantation). Seventy-two hours after NSC implantation, mouse brains were harvested and subjected to immunohistochemical analysis for the expression of the adenoviral early gene E1A (Figure 6, B, F, and J) and the late hexon gene (Figure 6, C, G, and K). Figure 6 illustrates viral hand-off by the presence of newly infected tumor cells as designated by positive staining for E1A. Tumor cells transitioning from an early to late phase of infection are represented by E1A hexon+ markers (Figure 6, J, K, and L, white arrow). Cells that are hexon+ denote a late phase of infection (Figure 6 J, K, and L, gray arrowhead). Based on these data, we conclude that OV-loaded HB1.F3.CD cells implanted in the contralateral hemisphere of glioma-bearing mice can home to, amplify, and hand off their therapeutic payload at distant tumor sites.

In Vivo Evaluation of the Therapeutic Efficacy of Distantly Delivered NSC-Based Oncolytic Virotherapy

Finally, we set out to evaluate the therapeutic potential of the OV-loaded HB1.F3.CD cell carrier when delivered at a site distant
from the primary tumor mass. We stereotactically implanted $5 \times 10^5$ OV-loaded (50 IU/cell) HB1.F3.CD cells in the left hemisphere of control mice (no tumor) or mice bearing U87 xenografts in the contralateral hemisphere. In the U87 model, median survival of the group treated with CRAd-S-pk7–loaded HB1.F3.CD cells was 33 days, which represented an 18% increase in median survival over the OV group (median survival = 28 days; $P < .001$) and a 16% increase over the HB1.F3.CD–treated group (median survival = 28.5 days; $P < .001$) (Figure 7A). In the GBM43FL model, an identical injection protocol was used as described in Figure 7A. The median survival was 22 days ($P = .01$) for the mice treated with HB1.F3.CD alone, 24 days ($P = .03$) for the CRAd-S-pk7–treated group, and 28.5 days for the HB1.F3.CD–loaded CRAd-S-pk7 group (Figure 7B). These data indicate that NSC carrier-based OV therapy can be effective even when implanted in the contralateral hemisphere of the targeted glioma xenograft.

**Discussion**

NSC carrier-based oncolytic virotherapy holds great promise as an alternative and complimentary treatment modality for glioblastoma.

In this study, we sought to evaluate the relative efficacy of two immortalized NSC lines as a cell carrier for antiglioma oncolytic virotherapy and observed that 1) the HB1.F3.CD cells were more efficient in supporting CRAd-S-pk7 OV replication as well as killing glioma cells both in vitro and in vivo compared with ReNcells; 2) the antiglioma activity of OV-loaded HB1.F3.CD cells was effective against clinically relevant human-derived glioma models as well as a GSC-enriched xenograft model; 3) OV-loaded HB1.F3.CD cells can effectively migrate to the contralateral hemisphere and hand off their therapeutic payload of oncolytic viruses to targeted glioma cells; 4) in vivo distribution and migratory kinetics of the OV-loaded HB1.F3.CD cells can be monitored by MRI and are reported in detail; and 5) distance delivery of OV-loaded HB1.F3.CD cells can prolong median survival in orthotopic mouse models with human glioma xenografts. These results along with our previously published data (8,9,15,16) argue in favor of NSC-based cell carriers for the targeted delivery of oncolytic viruses against human glioma. We believe that the methods and protocols as well as preclinical data generated during this study will permit us to bridge the gap between preclinical animal studies and human clinical trials and will lead to the development of human clinical trial protocols in the future.

Despite these promising results, certain important points of emphasis deserve further attention. First, we witnessed variability...
in the therapeutic efficacy of NSC-based oncolytic virotherapy between different glioma models. The models in our study were selected to represent the interpatient and intratumor heterogeneity widely observed among GBMs. On a molecular level, the GBM43 xenograft model expresses p53 mutant and PTEN wild-type, is negative for epidermal growth factor receptor, and expresses elevated levels of the chemo-resistance gene O6-methylguanine-DNA methyltransferase (MGMT) (12,17). The GBM12 model expresses p53 null status and wild-type PTEN (12,17) but expresses epidermal growth factor receptor. On the other hand, U87 contains wild-type p53, PTEN mutant, expresses low levels of wild-type epidermal growth factor receptor, and is MGMT negative (18). With respect to their pathological features, the U87 cell line forms very localized noninfiltrative tumors, whereas GBM43 tumors are mildly invasive, and GBM12 tumors are extremely infiltrative in the rodent brain. It has been well documented that the genetic phenotype of tumor cells is one of the key determining factors of antitumor activity of an oncolytic virus (19). For example, wild-type p53 status in tumor cells can facilitate an efficient killing of these tumor cells by OVs (11,12,20). This is consistent with the observation in our study that demonstrated the most pronounced efficacy in the p53 wild-type U87 glioma model.

Furthermore, the data presented in this report convincingly demonstrate that within 24 hours after implantation, OV-loaded HB1.F3.CD cells are capable of migrating 4 to 6 mm to the contralateral hemisphere and handing off therapeutic virus to targeted tumor cells in a rodent model of glioma. However, the smaller-sized brain makes it difficult to speculate on the migratory capacity of HB1.F3.CD cells in a larger-sized human brain with glioma. Fortunately, to mitigate the uncertainty behind this possible pitfall, we were able to demonstrate and optimize a valuable protocol where NSCs can be labeled with MPIOs and subsequently tracked by MRI imaging; therefore this technique may help serve as a key tool for the dynamic optimization of NSC delivery protocols in the clinical setting.

The fate of the NSC-based cell carrier is critical for maintaining tumor pathotropism (21). A number of recent studies reported that pluripotency and the differentiation status of the NSC are altered upon viral infection (22). Our data indicate that upon loading with the OV CRAd-S-pk7, the NSC-specific markers (nestin, Sox2 and Oct4) or different differentiation markers (Gal C, beta III tubulin, or GFAP) at the mRNA level remain unchanged. Moreover, the fate of the transplanted NSC in the mammalian brain has been shown to be influenced by the extrinsic factors that can drive their differentiation into either neurons or glia (23). The diseased brain seems to create an environment that can stimulate endogenous or exogenous stem cells to differentiate into specific cells types. However, our data, along with other published data, indicate that therapeutic stem cells implanted into the brains of tumor-bearing mice remain in an undifferentiated state (Figure 5) (24,25). These results strongly suggest that the tumor microenvironment may be deficient in factors necessary for stem cell differentiation.

Last, NSC allorejection could be detrimental to the success of NSC-based antiglioma oncolytic virotherapy in the clinic. Despite

**Figure 5.** In vivo differentiation of CRAd-S-pk7–loaded HB1.F3.CD cells. To follow the differentiation status of oncolytic adenovirus–loaded HB1.F3.CD cells implanted into the brains of mice, HB1.F3.CD-GFP+ loaded with CRAd-S-pk7 (50 IU per cell) were implanted into the contralateral hemisphere of mice bearing U87 glioma xenografts. Mice were killed 24 and 72 hours after neural stem cell (NSC) implantation, and brains were prepared for immunohistochemical analysis. A Hematoxylin and eosin staining of the migratory path of NSCs from the injection site represented by the (*) to the tumor site (A1). Magnified views of the migration path: injection site (A2), center of migration path (A3), end of migration site or tumor (A4). B Hematoxylin and eosin staining was confirmed by 4',6-diamidino-2-phenylindole (DAPI).
Figure 6. Neural stem cells (NSCs) hand off and expand oncolytic adenoviral virotherapy (OV) therapeutic payload at distant tumor foci in vivo. OV-loaded HB1.F3.CD cells were implanted in the contralateral hemisphere of mice bearing U87 xenograft tumors. Mice were killed 72 hours after NSC implantation, and brains were preserved and prepared for immunohistochemical analysis. A–D) stained with 4′,6-diamidino-2-phenylindole (DAPI) only. Scale bars: A–D = 200 μm, E–H = 100 μm I–L = 50 μm. B, F, J) Early stages of viral replication represented by the positive staining for E1A (gray arrowhead) inside the tumor border (dotted line). C, G, K) Intermediate stages of viral replication denoted by the costaining of E1A and hexon. D, H, L) (white arrow) Hexon positive staining represents the late phases of viral infection (white arrowhead). Sizing of scale bars: Figure 6 A-D (200 μm); Figure 6 E-H (100 μm); Figure 6 I-L (50 μm).

Figure 7. Contralateral delivery of oncolytic adenoviral virotherapy (OV)-loaded HB1.F3.CD cells show therapeutic efficacy in a mouse model of glioma. To examine distance delivery, either 5×10^3 U87 or GBM43FL cells were implanted into the right hemisphere of nude mice (n = 7 per group). After 3 days, mice were treated with an injection of either phosphate-buffered saline (PBS), 5×10^6 HB1.F3.CD cells, 5×10^6 HB1.F3.CD cells loaded with 50 IU per cell of CRAd-S-pk7, or 2.5×10^7 IU of CRAd-S-pk7 into the contralateral left hemisphere of the brain. A) Overall survival of U87MG bearing mice. B) Overall survival of GBM43FL bearing mice. CRAd-S-pk7-loaded HB1.F3.CD extended the efficacy of OV alone in both U87MG and GBM43FL models (P < .001, P = .03, respectively). P represents P value 1 in this table where PBS treated group served as reference and P represents P value 2 where CRAd-S-pk7+HB1.F3.CD serve as reference. Survival curves were obtained by the Kaplan–Meier method, and overall survival time was compared between groups using the log-rank test. All statistical tests were two-sided. *P < .05, **P < .01, ***P < .001.
the poor immunogenic characteristics of NSCs, in vitro NSC allorejection by peripheral lymphocytes has been reported (14,26). The ideal solution to this possible pitfall would be using autologous NSCs, but the currently available technology for isolating and expanding autologous NSCs is in its infancy. With recent developments in the field of induced pluripotent stem cells, one might imagine a future in which medical centers can offer highly customized, patient-focused approaches to NSC-based antiangioma therapy by using personalized induced pluripotent stem cells or induced NSCs (27) catered toward each patient with unique genetic backgrounds (27–29).

In conclusion, our study shows that NSC-based cell carriers can effectively deliver antiangioma OV to distant tumor sites, release the therapeutic payload at the target sites, and increase median survival in a diverse range of orthotopic human glioma xenograft models that stand to recapitulate the heterogeneity of the human disease. The importance of this point cannot be emphasized enough, because after all, the name “glioblastoma multiforme” does indeed mean “to have many forms, of multiple shapes,” which perfectly captures the essence of the multifaceted nature of the disease and the challenge it presents to developing an effective therapy. Thus, any novel antiangioma therapy must stand to address this crucial issue. In this study, we have demonstrated that the NSC OV platform has the ability to extend survival in a multitude of invasive models of human glioma and target the therapy-resistant and disease-reinitiating GSC population, thus fulfilling two critical considerations for successful clinical translation, and may serve as a future therapy that can complement the existing standard of care for glioblastoma.

References

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